WE CLAIM:

    1. An agonist analogue of amylin having the amino acid sequence depicted in Figure 3 wherein

        A_1 is Lys, Ala, Ser or hydrogen,

        B_1 is Ala, Ser or Thr;

        C_1 is Val, Leu or Ile;

        D_1 is His or Arg;

        E_1 is Ser or Thr;

        F_1 is Ser, Thr, Gln or Asn;

        G_1 is Asn, Gln or His;

        H_1 is Phe, Leu or Tyr;

        I_1 is Ala or Pro;

        J_1 is Ile, Val, Ala or Leu;

        K_1 is Ser, Pro, Leu, Ile or Thr;

        L_1 is Ser, Pro or Thr;

        M_1 is Asn, Asp, Gln or Asn; X and Y are independently selected residues having side chains which are chemically bonded to each other to form an intramolecular linkage; and Z is amino, alkylamino, dialkylamino, cycloalkylamino, arylamino, aralkylamino, alkyloxy, aryloxy or aralkyloxy; and provided that when

(a) A_1 is Lys, B_1 is Ala, C_1 is Val, D_1 is His, E_1 is Ser, F_1 is Ser, G_1 is Asn, H_1 is Phe, I_1 is Ala, J_1 is Ile, K_1 is Ser, L_1 is Ser, and M_1 is Asn;

(b) A_1 is Lys, B_1 is Ala, C_1 is Ile, D_1 is Arg, E_1 is Ser, F_1 is Ser, G_1 is Asn, H_1 is Leu, I_1 is Ala, J_1 is Ile, K_1 is Ser, L_1 is Pro, and M_1 is Asn;

(c) A_1 is Lys, B_1 is Ala, C_1 is Val, D_1 is Arg, E_1 is Thr, F_1 is Ser, G_1 is Asn, H_1 is Leu, I_1 is Ala, J_1 is Ile, K_1 is Ser, L_1 is Pro, and M_1 is Asn;

(d) A_1 is Lys, B_1 is Ala, C_1 is Val, D_1 is Arg, E_1 is Ser, F_1 is Ser, G_1 is Asn, H_1 is Leu, I_1 is Pro, J_1 is Val, K_1 is Pro, L_1 is Pro, and M_1 is Asn;

(e) A_1 is Lys, B_1 is Ala, C_1 is Val, D_1 is His, E_1 is Ser, F_1 is Asn, G_1 is Asn, H_1 is Leu, I_1 is Pro, J_1 is Val, K_1 is Ser, L_1 is Pro, and M_1 is Asn; or

(f) A_1 is Lys, B_1 is Thr, C_1 is Val, D_1 is Arg, E_1 is Ser, F_1 is Ser, G_1 is His, H_1 is Leu, I_1 is Ala, J_1 is Ala, K_1 is Leu, L_1 is Pro, and M_1 is Asp;

then one or more of any of A_1 to M_1 is not an L-amino acid and Z is not amino.